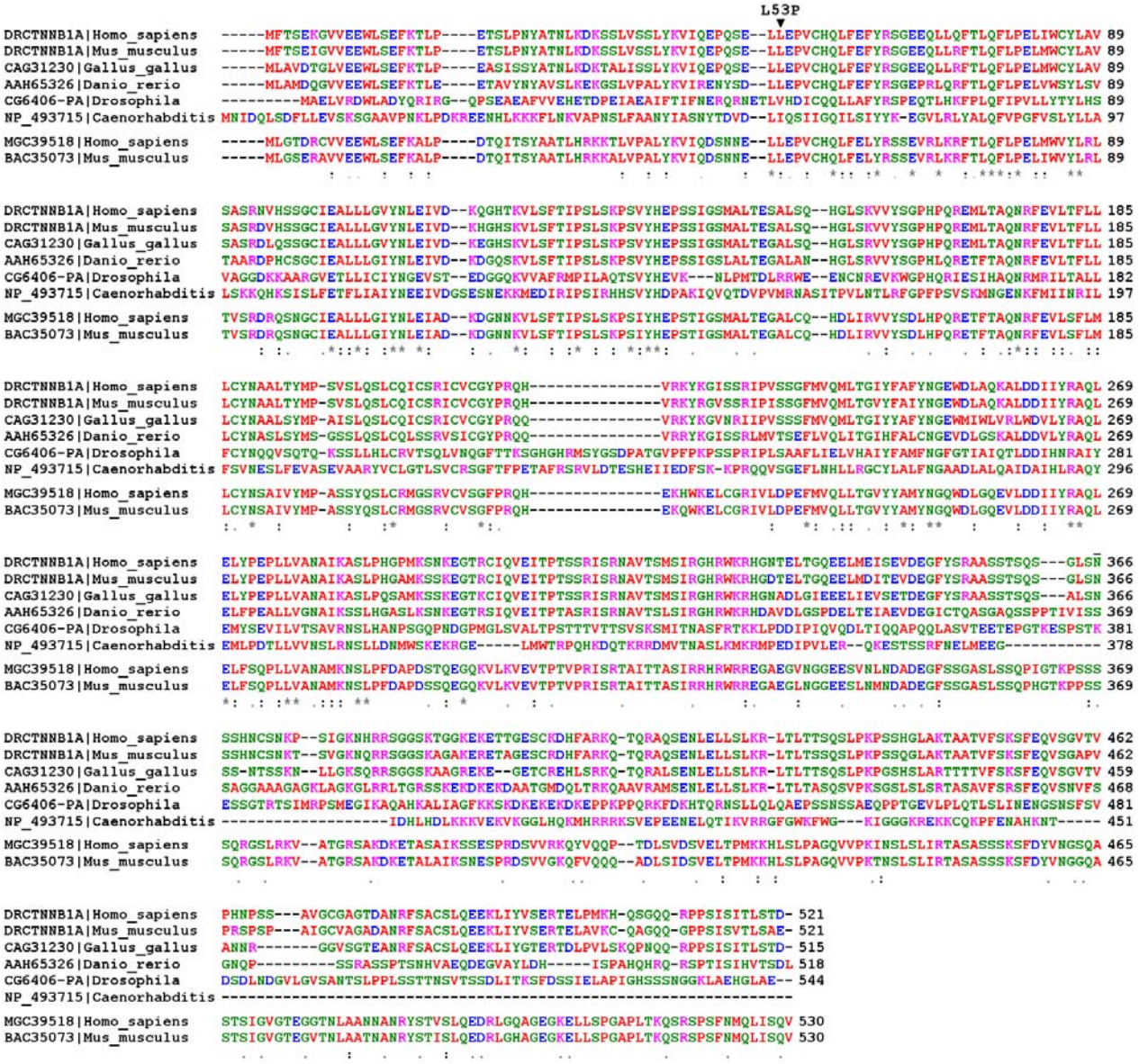
Supplementary figure 2: Multiple sequence alignment of hyccin.

Multiple sequence alignment of hyccin with 5 orthologues using CLUSTALW. The alignment of hyccin with a human paralogue and its murine orthologue is also shown. Residues conserved in all sequences are indicated by * symbol; Conserved and semi-conserved substitutions are indicated by " : " and " . " symbols, respectively. The L53P mutation is indicated with an arrow.